

SEQUENCE LISTING

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<120> INTERFERON-ALPHA INDUCED GENE

<130> 046658/298641

<160> 2

<170> PatentIn version 3.1

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<211> 6045
<212> DNA
<213> Homo sapiens

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<221> CDS
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Val Leu Leu	Ser Gln Lys Phe Asp	Asn Arg Gly Arg Lys	Met Ser	
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ctt aat gcc	tat gca ctg gat ttc	tac aaa cat ggt tcc	ttg ata	5198
Leu Asn Ala	Tyr Ala Leu Asp Phe	Tyr Lys His Gly Ser	Leu Ile	
1640	1645	1650		
gga tta gtc	cag gat aac agg atg	aat gaa gga gat gct	tat tat	5243

Gly	Leu	Val	Gln	Asp	Asn	Arg	Met	Asn	Glu	Gly	Asp	Ala	Tyr	Tyr		
		1655					1660					1665				
ttg	ttg	aag	gat	ttt	gca	ctc	acc	att	aaa	tct	atc	agt	gtt	tcc		5288
Leu	Leu	Lys	Asp	Phe	Ala	Leu	Thr	Ile	Lys	Ser	Ile	Ser	Val	Ser		
		1670					1675					1680				
ttg	cgt	gag	cta	tgt	gaa	aat	gaa	gac	gac	aac	gtt	gtc	tta	gcc		5333
Leu	Arg	Glu	Leu	Cys	Glu	Asn	Glu	Asp	Asp	Asn	Val	Val	Leu	Ala		
		1685					1690					1695				
ttt	gaa	caa	ctg	agt	aca	act	ttt	tgg	gaa	aag	tta	aac	aaa	gtc		5378
Phe	Glu	Gln	Leu	Ser	Thr	Thr	Phe	Trp	Glu	Lys	Leu	Asn	Lys	Val		
		1700					1705					1710				
taa	aaacaaagtc	tatgcaaacc	acttaaaaaat	aattccatag	tagttttttca											5431
ggtcacg	ttt	ttgattc	tta	tgcttcttgc	cagaaataca	ttatgataaa	gtggaaatac									5491
attacgatga	agtggaaaga	gcaaacactt	tggaatcaaa	cagagttgca	atcaaacctg											5551
ccatgttctg	tcatgaatac	tcacaaatta	tttagtatac	ctgaatcttg	gttttctttt											5611
ataactgagt	aataatgggt	acatctcagg	tagtttgagg	attgactaaa	aaaatgcgag											5671
aatgttgtat	gtgactgaat	aacaattttt	actctgcgaa	gccaaaagtaa	atataatatt											5731
atcagtaact	ttatccccag	tgtcagtatt	tataaaatgt	ttattaaggc	tagaaaaaat											5791
gaatacaata	tcctgaaggt	gaaatatatt	ctcttcaatt	agcataaata	tgattttacat											5851
aagttagcta	tacagctatt	gagatagtag	tttctagtaa	acttaaaacta	cttttttaaac											5911
atacattttg	tgatgattta	acaaaaatat	agagaatgat	ttgcttttatt	gtaattgtat											5971
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aaaaaaaaaa	aaaa															6045

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 <211> 1712
 <212> PRT
 <213> Homo sapiens

<400> 2

Met	Glu	Arg	Asn	Val	Leu	Thr	Thr	Phe	Ser	Gln	Glu	Met	Ser	Gln	Leu	
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Ile	Leu	Asn	Glu	Met	Pro	Lys	Ala	Glu	Tyr	Ser	Ser	Leu	Phe	Asn	Asp	
		20						25					30			
Phe	Val	Glu	Ser	Glu	Phe	Phe	Leu	Ile	Asp	Gly	Asp	Ser	Leu	Leu	Ile	
		35					40					45				
Thr	Cys	Ile	Cys	Glu	Ile	Ser	Phe	Lys	Pro	Gly	Gln	Asn	Leu	His	Phe	
	50					55					60					
Phe	Tyr	Leu	Val	Glu	Arg	Tyr	Leu	Val	Asp	Leu	Ile	Ser	Lys	Gly	Gly	
65				70					75					80		
Gln	Phe	Thr	Ile	Val	Phe	Phe	Lys	Asp	Ala	Glu	Tyr	Ala	Tyr	Phe	Asn	
			85						90					95		
Phe	Pro	Glu	Leu	Leu	Ser	Leu	Arg	Thr	Ala	Leu	Ile	Leu	His	Leu	Gln	

100					105					110					
Lys	Asn	Thr	Thr	Ile	Asp	Val	Arg	Thr	Thr	Phe	Ser	Arg	Cys	Leu	Ser
		115					120					125			
Lys	Glu	Trp	Gly	Ser	Phe	Leu	Glu	Glu	Ser	Tyr	Pro	Tyr	Phe	Leu	Ile
	130					135					140				
Val	Ala	Asp	Glu	Gly	Leu	Asn	Asp	Leu	Gln	Thr	Gln	Leu	Phe	Asn	Phe
145					150					155					160
Leu	Ile	Ile	His	Ser	Trp	Ala	Arg	Lys	Val	Asn	Val	Val	Leu	Ser	Ser
			165						170					175	
Gly	Gln	Glu	Ser	Asp	Val	Leu	Cys	Leu	Tyr	Ala	Tyr	Leu	Leu	Pro	Ser
			180					185					190		
Met	Tyr	Arg	His	Gln	Ile	Phe	Ser	Trp	Lys	Asn	Lys	Gln	Asn	Ile	Lys
		195					200					205			
Asp	Ala	Tyr	Thr	Thr	Leu	Leu	Asn	Gln	Leu	Glu	Arg	Phe	Lys	Leu	Ser
	210					215					220				
Ala	Leu	Ala	Pro	Leu	Phe	Gly	Ser	Leu	Lys	Trp	Asn	Asn	Ile	Thr	Glu
225					230					235					240
Glu	Ala	His	Lys	Thr	Val	Ser	Leu	Leu	Thr	Gln	Val	Trp	Pro	Glu	Gly
			245						250					255	
Ser	Asp	Ile	Arg	Arg	Val	Phe	Cys	Val	Thr	Ser	Cys	Ser	Leu	Ser	Leu
		260					265						270		
Arg	Met	Tyr	His	Arg	Phe	Leu	Gly	Asn	Arg	Glu	Pro	Ser	Ser	Gly	Gln
		275					280					285			
Glu	Thr	Glu	Ile	Gln	Gln	Val	Asn	Ser	Asn	Cys	Leu	Thr	Leu	Gln	Glu
	290					295					300				
Met	Glu	Asp	Leu	Cys	Lys	Leu	His	Cys	Leu	Thr	Val	Val	Phe	Leu	Leu
305				310					315						320
His	Leu	Pro	Leu	Ser	Gln	Arg	Ala	Cys	Ala	Arg	Val	Ile	Thr	Ser	His
			325					330					335		
Trp	Ala	Glu	Asp	Met	Lys	Pro	Leu	Leu	Gln	Met	Lys	Lys	Trp	Cys	Glu
		340						345					350		
Tyr	Phe	Ile	Leu	Arg	Asn	Ile	His	Thr	Phe	Glu	Phe	Trp	Asn	Leu	Asn
	355					360					365				
Leu	Ile	His	Leu	Ser	Asp	Leu	Asn	Asp	Glu	Leu	Leu	Leu	Lys	Asn	Ile
	370					375					380				
Ala	Phe	Tyr	Tyr	Glu	Asn	Glu	Asn	Val	Lys	Gly	Leu	His	Leu	Asn	Leu
385				390						395					400
Gly	Asp	Thr	Ile	Met	Lys	Asp	Tyr	Glu	Tyr	Leu	Trp	Asn	Thr	Val	Ser

705		710		715		720
Val Lys Val Lys Lys Arg Asn Lys Tyr Ser Ile Gly Ile Gly Pro Ala						
		725		730		735
Arg Phe Gln Leu Gln Tyr Met Gly His Tyr Leu Ile Arg Asp Glu Arg						
		740		745		750
Lys Asp Pro Asp Pro Arg Val Gln Asp Phe Ile Pro Asp Thr Trp Gln						
		755		760		765
Arg Glu Leu Leu Asp Val Val Asp Lys Asn Glu Ser Ala Val Ile Val						
		770		775		780
Ala Pro Thr Ser Ser Gly Lys Thr Tyr Ala Ser Tyr Tyr Cys Met Glu						
		785		790		800
Lys Val Leu Lys Glu Ser Asp Asp Gly Val Val Val Tyr Val Ala Pro						
		805		810		815
Thr Lys Ala Leu Val Asn Gln Val Ala Ala Thr Val Gln Asn Arg Phe						
		820		825		830
Thr Lys Asn Leu Pro Ser Gly Glu Val Leu Cys Gly Val Phe Thr Arg						
		835		840		845
Glu Tyr Arg His Asp Ala Leu Asn Cys Gln Val Leu Ile Thr Val Pro						
		850		855		860
Ala Cys Phe Glu Ile Leu Leu Leu Ala Pro His Arg Gln Asn Trp Val						
		865		870		875
Lys Lys Ile Arg Tyr Val Ile Phe Asp Glu Val His Cys Leu Gly Gly						
		885		890		895
Glu Ile Gly Ala Glu Ile Trp Glu His Leu Leu Val Met Ile Arg Cys						
	900		905		910	
Pro Phe Leu Ala Leu Ser Ala Thr Ile Ser Asn Pro Glu His Leu Thr						
	915		920		925	
Glu Trp Leu Gln Ser Val Lys Trp Tyr Trp Lys Gln Glu Asp Lys Ile						
	930		935		940	
Ile Glu Asn Asn Thr Ala Ser Lys Arg His Val Gly Arg Gln Ala Gly						
	945		950		955	960
Phe Pro Lys Asp Tyr Leu Gln Val Lys Gln Ser Tyr Lys Val Arg Leu						
	965		970		975	
Val Leu Tyr Gly Glu Arg Tyr Asn Asp Leu Glu Lys His Val Cys Ser						
	980		985		990	
Ile Lys His Gly Asp Ile His Phe Asp His Phe His Pro Cys Ala Ala						
	995		1000		1005	

Leu	Thr	Thr	Asp	His	Ile	Glu	Arg	Tyr	Gly	Phe	Pro	Pro	Asp	Leu
1010						1015					1020			
Thr	Leu	Ser	Pro	Arg	Glu	Ser	Ile	Gln	Leu	Tyr	Asp	Ala	Met	Phe
1025						1030					1035			
Gln	Ile	Trp	Lys	Ser	Trp	Pro	Arg	Ala	Gln	Glu	Leu	Cys	Pro	Glu
1040						1045					1050			
Asn	Phe	Ile	His	Phe	Asn	Asn	Lys	Leu	Val	Ile	Lys	Lys	Met	Asp
1055						1060					1065			
Ala	Arg	Lys	Tyr	Glu	Glu	Ser	Leu	Lys	Ala	Glu	Leu	Thr	Ser	Trp
1070						1075					1080			
Ile	Lys	Asn	Gly	Asn	Val	Glu	Gln	Ala	Arg	Met	Val	Leu	Gln	Asn
1085						1090					1095			
Leu	Ser	Pro	Glu	Ala	Asp	Leu	Ser	Pro	Glu	Asn	Met	Ile	Thr	Met
1100						1105					1110			
Phe	Pro	Leu	Leu	Val	Glu	Lys	Leu	Arg	Lys	Met	Glu	Lys	Leu	Pro
1115						1120					1125			
Ala	Leu	Phe	Phe	Leu	Phe	Lys	Leu	Gly	Ala	Val	Glu	Asn	Ala	Ala
1130						1135					1140			
Glu	Ser	Val	Ser	Thr	Phe	Leu	Lys	Lys	Lys	Gln	Glu	Thr	Lys	Arg
1145						1150					1155			
Pro	Pro	Lys	Ala	Asp	Lys	Glu	Ala	His	Val	Met	Ala	Asn	Lys	Leu
1160						1165					1170			
Arg	Lys	Val	Lys	Lys	Ser	Ile	Glu	Lys	Gln	Lys	Ile	Ile	Asp	Glu
1175						1180					1185			
Lys	Ser	Gln	Lys	Lys	Thr	Arg	Asn	Val	Asp	Gln	Ser	Leu	Ile	His
1190						1195					1200			
Glu	Ala	Glu	His	Asp	Asn	Leu	Val	Lys	Cys	Leu	Glu	Lys	Asn	Leu
1205						1210					1215			
Glu	Ile	Pro	Gln	Asp	Cys	Thr	Tyr	Ala	Asp	Gln	Lys	Ala	Val	Asp
1220						1225					1230			
Thr	Glu	Thr	Leu	Gln	Lys	Val	Phe	Gly	Arg	Val	Lys	Phe	Glu	Arg
1235						1240					1245			
Lys	Gly	Glu	Glu	Leu	Lys	Ala	Leu	Ala	Glu	Arg	Gly	Ile	Gly	Tyr
1250						1255					1260			
His	His	Ser	Ala	Met	Ser	Phe	Lys	Glu	Lys	Gln	Leu	Val	Glu	Ile
1265						1270					1275			
Leu	Phe	Arg	Lys	Gly	Tyr	Leu	Arg	Val	Val	Thr	Ala	Thr	Gly	Thr
1280						1285					1290			

Leu Ala 1295	Leu Gly Val Asn Met 1300	Pro Cys Lys Ser Val 1305	Val Phe Ala
Gln Asn 1310	Ser Val Tyr Leu Asp 1315	Ala Leu Asn Tyr Arg 1320	Gln Met Ser
Gly Arg 1325	Ala Gly Arg Arg Gly 1330	Gln Asp Leu Met Gly 1335	Asp Val Tyr
Phe Phe 1340	Asp Ile Pro Phe Pro 1345	Lys Ile Gly Lys Leu 1350	Ile Lys Ser
Asn Val 1355	Pro Glu Leu Arg Gly 1360	His Phe Pro Leu Ser 1365	Ile Thr Leu
Val Leu 1370	Arg Leu Met Leu Leu 1375	Ala Ser Lys Gly Asp 1380	Asp Pro Glu
Asp Thr 1385	Lys Ala Lys Val Leu 1390	Ser Val Leu Lys His 1395	Ser Leu Leu
Ser Phe 1400	Lys Gln Pro Arg Val 1405	Met Asp Met Leu Lys 1410	Leu Tyr Phe
Leu Phe 1415	Ser Leu Gln Phe Leu 1420	Val Lys Glu Gly Tyr 1425	Leu Asp Gln
Glu Gly 1430	Asn Pro Met Gly Phe 1435	Ala Gly Leu Val Ser 1440	His Leu His
Tyr His 1445	Glu Pro Ser Asn Leu 1450	Val Phe Val Ser Phe 1455	Leu Val Asn
Gly Leu 1460	Phe His Asp Leu Cys 1465	Gln Pro Thr Arg Lys 1470	Gly Ser Lys
His Phe 1475	Ser Gln Asp Val Met 1480	Glu Lys Leu Val Leu 1485	Val Leu Ala
His Leu 1490	Phe Gly Arg Arg Tyr 1495	Phe Pro Pro Lys Phe 1500	Gln Asp Ala
His Phe 1505	Glu Phe Tyr Gln Ser 1510	Lys Val Phe Leu Asp 1515	Asp Leu Pro
Glu Asp 1520	Phe Ser Asp Ala Leu 1525	Asp Glu Tyr Asn Met 1530	Lys Ile Met
Glu Asp 1535	Phe Thr Thr Phe Leu 1540	Arg Ile Val Ser Lys 1545	Leu Ala Asp
Met Asn 1550	Gln Glu Tyr Gln Leu 1555	Pro Leu Ser Lys Ile 1560	Lys Phe Thr
Gly Lys 1565	Glu Cys Glu Asp Ser 1570	Gln Leu Val Ser His 1575	Leu Met Ser

Cys	Lys	Glu	Gly	Arg	Val	Ala	Ile	Ser	Pro	Phe	Val	Cys	Leu	Ser
1580						1585					1590			
Gly	Asn	Phe	Asp	Asp	Asp	Leu	Leu	Arg	Leu	Glu	Thr	Pro	Asn	His
1595						1600					1605			
Val	Thr	Leu	Gly	Thr	Ile	Gly	Val	Asn	Arg	Ser	Gln	Ala	Pro	Val
1610						1615					1620			
Leu	Leu	Ser	Gln	Lys	Phe	Asp	Asn	Arg	Gly	Arg	Lys	Met	Ser	Leu
1625						1630					1635			
Asn	Ala	Tyr	Ala	Leu	Asp	Phe	Tyr	Lys	His	Gly	Ser	Leu	Ile	Gly
1640						1645					1650			
Leu	Val	Gln	Asp	Asn	Arg	Met	Asn	Glu	Gly	Asp	Ala	Tyr	Tyr	Leu
1655						1660					1665			
Leu	Lys	Asp	Phe	Ala	Leu	Thr	Ile	Lys	Ser	Ile	Ser	Val	Ser	Leu
1670						1675					1680			
Arg	Glu	Leu	Cys	Glu	Asn	Glu	Asp	Asp	Asn	Val	Val	Leu	Ala	Phe
1685						1690					1695			
Glu	Gln	Leu	Ser	Thr	Thr	Phe	Trp	Glu	Lys	Leu	Asn	Lys	Val	
1700						1705					1710			